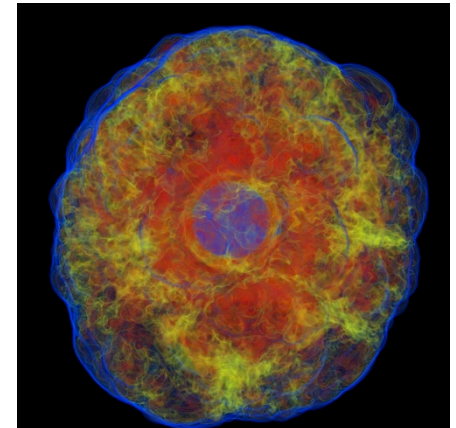
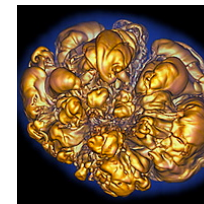
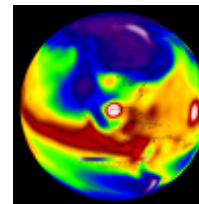
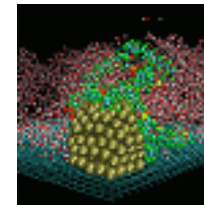
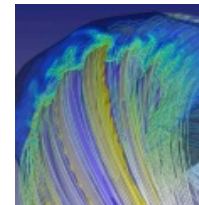
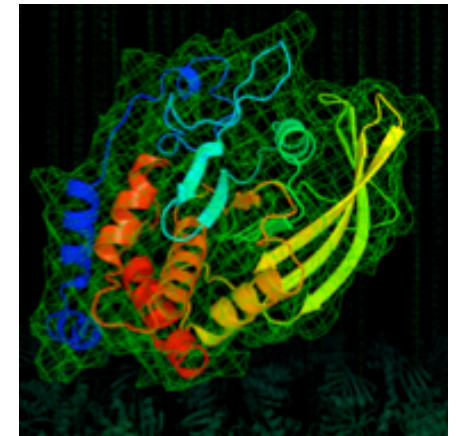
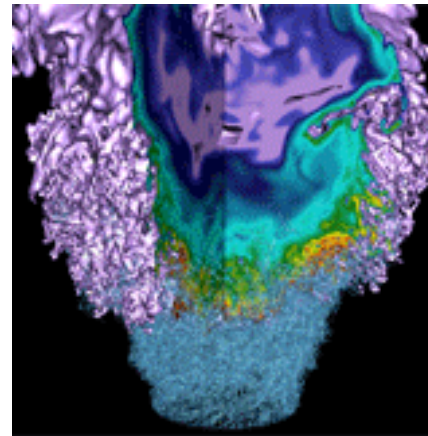


# Computing Environment



**Scott French**  
**NERSC User Services Group**

**New User Training**  
**August 13, 2015**

# Node Types

- **Login nodes**
  - Shared with other users
  - Code compilation, job preparation and submission
- **MOM nodes (Cray machines)**
  - Shared with other users
  - Where batch script executes
  - Will go away when we transition to SLURM
- **Compute nodes**
  - Not shared (except serial queues)

# Login Node Configuration

- **Edison**
  - Twelve nodes
    - 16 cores, 2.0 GHz Intel Sandy Bridge, 512 GB
- **Hopper**
  - Eight nodes
    - 16 cores, 2.4 GHz AMD Opteron, 128GB
  - Four nodes
    - 32 cores, 2.0 GHz AMD Opteron, 128GB
- **Genepool**
  - Four nodes
    - 8 cores, 2.3 GHz Intel Sandy Bridge, 32 GB
- **PDSF**
  - Three nodes
    - 16 cores, 2.6 GHz Intel Sandy Bridge, 125 GB

# Login Node Access

- **Connect (via ssh) to *load balancer***
  - % ssh **edison.nersc.gov**
  - % ssh **hopper.nersc.gov**
  - % ssh **genepool.nersc.gov**
  - % ssh **pdsf.nersc.gov**
- **Load balancer selects login node based on:**
  - Number of connections
  - Memory of previous connections from same IP

# Login Node Usage



- Login nodes are shared by many users, all the time
- Edit files, compile programs, submit batch jobs
- *Some* post-processing/data analysis
  - IDL, MATLAB, NCL, python, etc.
- *Some* file transfers
  - Use data transfer nodes for large/long-running transfers
- Please use discretion
  - *All* users get frustrated by sluggish interactive response

# Login Node Guidelines



- Use *no more* than 50% of available cores
- Use *no more* than 25% of available memory
- Limit use of parallel “make”  

```
% make -j 4 all
```
- NERSC will kill user processes if login nodes become unacceptably slow or unresponsive
- Terminate idle sessions of licensed software
  - IDL
  - MATLAB
  - Mathematica

# Shell Initialization Files

- **Standard dot files are maintained by NERSC**
  - `.bashrc`, `.profile`, `.cshrc`, `.login`, etc.
  - Symbolic links to read-only files
- **Personal dot files**
  - Aliases, environment variables, modules, etc.
  - Use `.ext` suffix (“`.ext` files”) `.bashrc.ext`, etc.
- **Broken? Use “`fixdots`” to start over**
  - Creates `$HOME/KeepDots.<timestamp>`
  - Restores all dot files to default state
  - If `PATH` corrupted:  
`/usr/common/usg/bin/fixdots`
- **Use NIM to change default login shell**

# NERSC Supported Software



- **NERSC provides a wide range of software**
  - Scientific Applications
    - VASP, Amber, NAMD, ABySS, ...
  - Compilers
    - Intel, GCC, PGI, Cray
  - Scripting Languages
    - perl, python, R - including common packages for each
  - Software Libraries (some maintained by Cray)
    - blas/lapack (MKL), boost, hdf5, netcdf, ...
  - Development utilities
    - git, mercurial, cmake, ...
  - Debuggers and Profilers
    - CrayPat, DDT, TotalView, gdb, MAP, darshan, IPM, VTune
  - Visualization
    - Visit, ParaView, VMD, ...
- **See complete list**  
<http://www.nersc.gov/users/software/>



# Software is Managed by Modules

- **Identify the software you need**

- Use the NERSC website

- <http://www.nersc.gov/users/software/>

- Use `module avail`

- *Lots* of output

- All module output goes to stderr, not stdout

- Each system has different modules!

- **Load the module**

```
% which idl
```

```
idl: Command not found.
```

```
% module load idl
```

```
% which idl
```

```
/usr/common/usg/idl/idl82/bin/idl
```

# Loading Modules

- **Different module for each version of software**
    - Syntax: <name>/<version>
    - Default provided if no <version> supplied
- ```
% module avail idl  
idl/7.1      idl/8.0      idl/8.2 (default)  
% module load idl/7.1
```
- **Load modules in every batch script**
    - Ensure correct run-time environment
    - Self-documenting for troubleshooting and reproducibility

# Other Useful Module Commands

**module unload <modulename>**

- Remove the module from your environment

**module swap <module1> <module2>**

- Unload one module and replace it with another

`module swap pgi gcc`

**module list**

- See what modules you have loaded right now

**module show <modulename>**

- See what the module actually does

**module help <modulename>**

- Get more information about the software

# Default Modules



- **When you login, many *default* modules are loaded automatically**
  - Usually foundational modules which are required to get proper function from the system
    - Build environment, required libraries and applications, batch environment
  - Use caution in unloading these
- **Swapping to functionally equivalent module may be OK**  
`hopper% module swap PrgEnv-pgi PrgEnv-gnu`
- **Each NERSC system has different default modules**

# Types of Modules

- **Applications**
  - VASP, amber, blast, ...
  - Usually only set `PATH`, `LD_LIBRARY_PATH`
- **Libraries**
  - Set `LD_LIBRARY_PATH`
    - but probably not on Crays
  - Set “helper” environment variable for building software
    - Header/include file search paths
    - Library search paths
    - Library names

```
% module load hdf5  
  
% mpicc mycode.f $HDF5
```

# Cray Programming Environment

- **Compiler specific**

PrgEnv-pgi, PrgEnv-intel, PrgEnv-cray, PrgEnv-gnu

- Intel is default on Edison, PGI is default on Hopper

- ***Meta-modules***

- Organize a set of modules

- Compiler (intel, pgi, cray, gnu)
- Libraries (including MPI) tuned for compiler

- **Swapping Programming Environments**

`module swap PrgEnv-pgi PrgEnv-intel`

- swaps compiler

- *no need to swap libraries!*

# Compiler Wrappers

- **On Hopper / Edison:**
  - Defined by `PrgEnv-*` modules
  - `ftn` (fortran), `cc` (C), `CC` (C++)
  - Provides include header and library search paths for MPI, common math libraries (e.g., Cray libsci), Cray system software
  - Provides consistent level of optimization across compilers
- **Seldom need native compilers!**

- **Provides different OS environments**
  - Often different third-party software
    - Some software packages have specific OS requirements
      - Possibly due to validation requirements
- **Used on Genepool, PDSF, and Carver (retires in September 2015)**
- **Transparent**
  - Default configuration for most users
  - Alternate configurations for some users
- **Details on website**

<http://www.nersc.gov/users/computational-systems/carver/user-environment/>

<http://www.nersc.gov/users/computational-systems/pdsf/software-and-tools/chos/>



# Resources

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<http://www.nersc.gov/users/software/nersc-user-environment/>

<http://www.nersc.gov/users/software/nersc-user-environment/modules/>

<http://www.nersc.gov/users/computational-systems/edison/programming>

<http://www.nersc.gov/users/computational-systems/hopper/programming/>

<http://www.nersc.gov/users/computational-systems/carver/programming/>



**Thank you.**